17-H PA. INT COOPERATION TREAT

|  | From the INTERNATIONAL BUREAU  |
|--|--|
| PCT  | To:  |
| NOTIFICATION OF ELECTION  (PCT Rule 61.2)  | Assistant Commissioner for Patents United States Patent and Trademark Office Box PCT Washington, D.C.20231 ÉTATS-UNIS D'AMÉRIQUE |
| Date of mailing (day/month/year)   |  |
| 21 January 2000 (21.01.00)   | in its capacity as elected Office  |
| International application No. PCT/US99/09761   | Applicant's or agent's file reference 44921-5004WO   |
| International filing date (day/month/year)   | Priority date (day/month/year)   |
| 05 May 1999 (05.05.99)   | 05 May 1998 (05.05.98)   |
| Applicant PRASHAR, Yatindra et al  |  |
| 1. The designated Office is hereby notified of its election made    X   in the demand filed with the International Preliminary   03 December 1 | Examining Authority on: 999 (03.12.99) ational Bureau on:  |

The International Bureau of WIPO 34, chemin des Colombettes 1211 Geneva 20, Switzerland Facsimile No.: (41-22) 740.14.35 Authorized officer

Olivia RANAIVOJAONA

Telephone No.: (41-22) 338.83.38

## RECEIVED

## TENT COOPERATION TRE

NOV 22 1999

MORGAN, LEWIS & BOCKIUS LLP PCT

NOTICE INFORMING THE APPLICANT OF THE COMMUNICATION OF THE INTERNATIONAL APPLICATION TO THE DESIGNATED OFFICES

(PCT Rule 47.1(c), first sentence)

#### From the INTERNATIONAL BUREAU

To:
ADLER, Reid, G.
Morgan, Lewis & Bockius LLP
1800 M Street, N.W.
Washington, DC 20036
ÉTATS-UNIS D'AMÉRIQUE

Date of mailing (day month year)

11 November 1999 (11.11.99)

Applicant's or agent's file reference

44921-5004WO

International application No. PCT US99 09761

International filing date (day month year) 05 May 1999 (05.05.99)

Priority date (day month year)

IMPORTANT NOTICE

05 May 1998 (05.05.98)

Applicant

GENE LOGIC, INC. et al.

 Notice is hereby given that the International Bureau has communicated, as provided in Article 20, the international application to the following designated Offices on the date indicated above as the date of mailing of this Notice: AU.EP.JP.US

In accordance with Rule 47.1(c), third sentence, those Offices will accept the present Notice as conclusive evidence that the communication of the international application has duly taken place on the date of mailing indicated above and no copy of the international application is required to be furnished by the applicant to the designated Office(s).

2. The following designated Offices have waived the requirement for such a colomunication at this time:

CA

The communication will be made to those Offices only upon their reduest. Furthermore, those Offices do not require the applicant to furnish a copy of the international application (Rule 49.1(a-bis)).

 Enclosed with this Notice is a copy of the international application as published by the International Bureau on 11 November 1999 (11.11.99) under No. WO 99 57130.

#### REMINDER REGARDING CHAPTER II (Article 31(2)(a) and Rule 54.2)

If the applicant wisnes to postpone entry into the national onase until 30 months (or later in some Offices) from the priority date, a demand for international preliminary examination must be filed with the competent International Preliminary Examining Authority before the expiration of 19 months from the priority date.

it is the applicant's sole responsibility to monitor the 19-month time limit

Note that only an applicant who is a national or resident of a PCT Contracting State which is bound by Chapter II has the right to file a demand for international preliminary examination.

#### REMINDER REGARDING ENTRY INTO THE NATIONAL PHASE (Article 22 or 39(1))

If the applicant wisnes to proceed with the international application in the **national phase**, ne must, within 20 months or 30 months, or later in some Offices, perform the acts referred to therein perfore each designated or elected Office.

For further important information on the time, imits and acts to be performed for entering the national phase, see the Annex to Form PCT B 301 (Notification of Receipt of Receipt Coby, and Volume), of the PCT Apolicant's Guide.

IEC 2 0 JUN 2000

# **PCT**

# INTERNATIONAL PRELIMINARY EXAMINATION REPORT

(PCT Articl 36 and Rule 70)

| Applicant's or agent's file reference 44921-5004WO  | FOR FURTHER ACTION   | ee Notification of Transmitta<br>reliminary Examination Report (Fo |  |  |  |  |
|---|--|--|--|--|--|--|
| International application No.   | International filing date (day/mo  | h/year) Priority date (day/mo                                      | nth/year)                                |  |  |  |
| PCT/US99/09761  | 05 MAY 1999  | 05 MAY 1998  | ,  |  |  |  |
| International Patent Classification (IPC) Please See Supplemental Sheet.  | L  | 33 MAT 1330  |  |  |  |  |
| Applicant GENE LOGIC, INC.  |  |  |  |  |  |  |
| Examining Authority and is  | ry examination report has b<br>transmitted to the applicant ac   | n prepared by this International ording to Article 36.             | onal Preliminary                         |  |  |  |
| 2. This REPORT consists of a total of sheets.   |  |  |  |  |  |  |
| been amended and are the  | panied by ANNEXES, i.e., sheets basis for this report and/or sheet ion 607 of the Administrative In tal of sheets. | containing rectifications made b                                   | rawings which have efore this Authority. |  |  |  |
| 3. This report contains indication  | s relating to the following iter   | :  |  |  |  |  |
| I X Basis of the repor  | t  |  |  |  |  |  |
| II Priority   |  |  |  |  |  |  |
| III Non-establishment of report with regard to novelty, inventive step or industrial applicability  |  |  |  |  |  |  |
| IV Lack of unity of invention   |  |  |  |  |  |  |
| V X Reasoned statement under Article 35(2) with regard to novelty, inventive step or industrial applicability, citations and explanations supporting such statement |  |  |  |  |  |  |
| VI Certain documents  | cited  |  |  |  |  |  |
| VII Certain defects in t  | he international application   |  |  |  |  |  |
| VIII Certain observation  | s on the international application   |  |  |  |  |  |
|   |  |  |  |  |  |  |
|   |  |  |  |  |  |  |
|   |  |  |  |  |  |  |
| Date of submission of the demand  | Date o   | completion of this report  |  |  |  |  |
| 03 DECEMBER 1999  | 18   | AY 2000  |  |  |  |  |
| Name and mailing address of the IPEA/   |  | ed officer   | llens/                                   |  |  |  |
| Commissioner of Patents and Tradem<br>Box PCT<br>Washington, D.C. 20231   | PA   | RICK NODAN   | elinox p                                 |  |  |  |
| Facsimile No. (703) 305-3230  | Teleph   | e No. (703) 308-0196   |  |  |  |  |

Form PCT/iPEA/409 (cover sheet) (July 1998)\*



## INTERNATIONAL PRELIMINARY EXAMINATION REPORT

| _           |                  |        |     |
|-------------|------------------|--------|-----|
|             | REC'D 20         | JUN 20 | 100 |
| Internation | ah application i | No.    |     |
| PCT/US      | 99/09761         |        | POT |

| L B     | asis f th         | ne rep rt                                       |                             |   |   |
|---------|-------------------|---|-----------------------------|---|---|
| 1. Witl | n regard to       | the elements of the interna                     | tional application:*        |   |   |
| x       | i .               | mational application as                         |                             |   |   |
| x       | the desc          | cription:                                       |                             |   |   |
|         | pages _           | 1-48  |                             |   | , as originally filed   |
|         | pages             | NONE  |                             |   | _, filed with the demand  |
|         | pages _           | NONE  | , filed                     | with the letter of  |   |
|         | the clair         | me.   |                             |   |   |
| X       | pages _           | 40.67   |                             |   | , as originally filed   |
|         | pages _           |   |                             |   | statement) under Article 19                                       |
|         | pages _           |   |                             |   | _ , filed with the demand   |
|         | pages _           | NONE  | , filed with the le         | tter of   |   |
|         | the draw          | vinge:  |                             |   |   |
| X       | the drav          | =   |                             |   | as originally filed   |
|         | pages _           | NONE  |                             |   | filed with the demand   |
|         | pages _           | NONE  | , filed w                   | rith the letter of  | _ , mos war als comans  |
|         |                   |   | •                           |   |   |
| X       | the sequ          | ence listing part of the d                      | escription:                 |   | _   |
|         |                   |   | 10                          |   | , as originally filed   |
|         | pages _           | NONE  | £1. 1                       | ide de lawer 6  | _ , filed with the demand   |
|         | pages _           | NUNE  | , filed w                   | ith the letter of   |   |
|         | the lang          | guage of publication of                         | the international applic    | es of international search (teation (under Rule 48.3(b)).  If international preliminary exa |   |
|         | or 55.3).         |   |                             |   |   |
|         | _                 | to any nucleotide and/o examination was carried | _                           |   | l application, the international                                  |
|         | containe          | ed in the international a                       | pplication in printed f     | orm.  |   |
|         | filed to          | gether with the internati                       | onal application in co      | mputer readable form.   |   |
|         | furnishe          | ed subsequently to this                         | Authority in written fo     | orm.  |   |
|         | furnishe          | ed subsequently to this                         | Authority in computer       | readable form.  |   |
|         |                   | tement that the subsequer                       |                             | equence listing does not go   | beyond the disclosure in the                                      |
|         | The state         |   | recorded in computer r      | eadable form is identical to the  | e writen sequence listing has                                     |
| 4. X    | The am            | nendments have resulted                         | in the cancellation of      | f:  |   |
|         | X t               | ne description, pages                           | N/A                         | <u>.</u>  |   |
|         | Ľ tł              | ne claims, Nos.                                 | N/A                         | <del></del>   |   |
|         | _ <u>  X</u>   tł | he drawings, sheets <del>/fig</del>             | N/A                         |   |   |
| 5. X    |                   |   |                             | s had not been made, since the  | ey have been considered to go                                     |
| in      | placement :       | sheets which have been fur                      | nished to the receiving Off | nental Box (Rule 70.2(c)).** fice in response to an invitation report since they do not con | under Article 14 are referred to<br>stain amendments (Rules 70.16 |
|         |                   | ment sheet containing suc                       | h amendments must be t      | eferred to under item 1 and i   | annexed to this report.   |





#### INTERNATIONAL PRELIMINARY EXAMINATION REPORT

International application No.

PCT/US99/09761

|    | citations and explanations suppo | rting such statem | ent                           |     |
|----|----------------------------------|-------------------|-------------------------------|-----|
| 1. | statement                        |                   |                               |     |
|    | Novelty (N)                      | Claims            | 1-33                          | YES |
|    |                                  | Claims            | NONE                          | NO  |
|    | Inventive Step (IS)              | Claims            | 3, 8-11, 14-15, 24-25, 28-33  | YES |
|    |                                  | Claims            | 1-2, 4-7, 12-13, 16-23, 26-27 | NO  |

V. Reasoned statement under Article 35(2) with regard to novelty, inventive step or industrial applicability;

Industrial Applicability (IA)

Claims 1-33

Claims NONE

NO

### 2. citations and explanations (Rule 70.7)

Claims 1, 2, 4-7, 12-13, 16-23 and 26-27 lack an inventive step under PCT Article 33(3) as being obvious over WO 97/05286 in view of Grossman et al.

The '286 patent teaches using gene expression from activated and non-activated T cells and comparing the effects of drugs on said activation levels for use in treating diseases or comparing drug use in treatment outcomes.

The claimed invention differs from the prior art teachings by the reciation of using said gene expression method of T cells in the diagnosis or treatment protocols of HIV. However, Grossman et al., teaches using gene expression in T cells during HIV as a tool to determine disease progression and therapy.

Therefore it would have been prima facie obvious to one of ordinary skill in the art at the time the invention was made to use the assay taught by the '286 patent to determine T cell activation in HIVas taught by Grossman et al.

Claims 3, 8-11, 24-25 and 28-33 meet the criteria set out in PCT Article 33(3), because the prior art does not teach or fairly suggest using B cells in the assay or assaying for genetic diseases or the specific primers.

| Claims | 1-33 | meet | the | criteria | set | out | under | <b>PCT</b> | <b>Articles</b> | 33(2) | and | 33(4). |
|--------|------|------|-----|----------|-----|-----|-------|------------|-----------------|-------|-----|--------|
|        |      | }    | NEV | V CITA   | TIC | NS  |       |            |                 |       |     |        |
| NONE   |      |      |     |          |     |     |       |            |                 |       |     |        |





### INTERNATIONAL PRELIMINARY EXAMINATION REPORT

International application No. PCT/US99/09761

| Supp | lemental | Box |
|------|----------|-----|
|------|----------|-----|

(To be used when the space in any of the preceding boxes is not sufficient)

Continuation of: Boxes I - VIII

Sheet 10

| CI | .ASSI | FIC | 'AT | OF | N٠ |
|----|-------|-----|-----|----|----|
|    |       |     |     |    |    |

The International Patent Classification (IPC) and/or the National classification are as listed below: IPC(7): C07H 21/04; C12N 15/00; C12Q 1/02, 1/68 and US Cl.: 435/6, 29, 320.1, 325; 536/24.3, 23.5

- I. BASIS OF REPORT:
- 5. (Some) amendments are considered to go beyond the disclosure as filed: NONE

## **PCT**





|   | JNDER THE PATENT COOPERATION TREATY (PCT)          |   |
|---|--|---|
| (51) International Patent Classification 6:   | (11) International Publication Number: WO 99/57130 |   |
| C07H 21/04, C12N 15/00, C12Q 1/02, 1/68   | A1   | (43) International Publication Date: 11 November 1999 (11.11.99)  |
| (21) International Application Number: PCT/US   | \$99/097   | 61 (81) Designated States: AU, CA, JP, US, European patent (AT, BE, CH, CY, DE, DK, ES, FI, FR, GB, GR, IE, IT, LU, MC,                                 |
| (22) International Filing Date: 5 May 1999  | (05.05.9   | 9) NL, PT, SE).   |
| (30) Priority Data:   |  | Published   |
| 60/084,329 5 May 1998 (05.05.98)  | τ  | With international search report.  Before the expiration of the time limit for amending the claims and to be republished in the event of the receipt of |
| (71) Applicant (for all designated States except US): GEN INC. [US/US]; 708 Quince Orchard Road, Gai MD 21046 (US). |  | C, amendments.  |
| (72) Inventors; and   |  |   |

(74) Agent: ADLER, Reid, G.; Morgan, Lewis & Bockius LLP, 1800 M Street, N.W., Washington, DC 20036 (US).

Ronan Street, New Haven, CT 06511 (US).

(75) Inventors/Applicants (for US only): PRASHAR, Yatindra [IN/US]; 9749 Clocktower Lane #304, Columbia, MD 21046 (US). WEISSMAN, Sherman [US/US]; 459 St.

(54) Title: A PROCESS TO STUDY CHANGES IN GENE EXPRESSION IN T LYMPHOCYTES

#### (57) Abstract

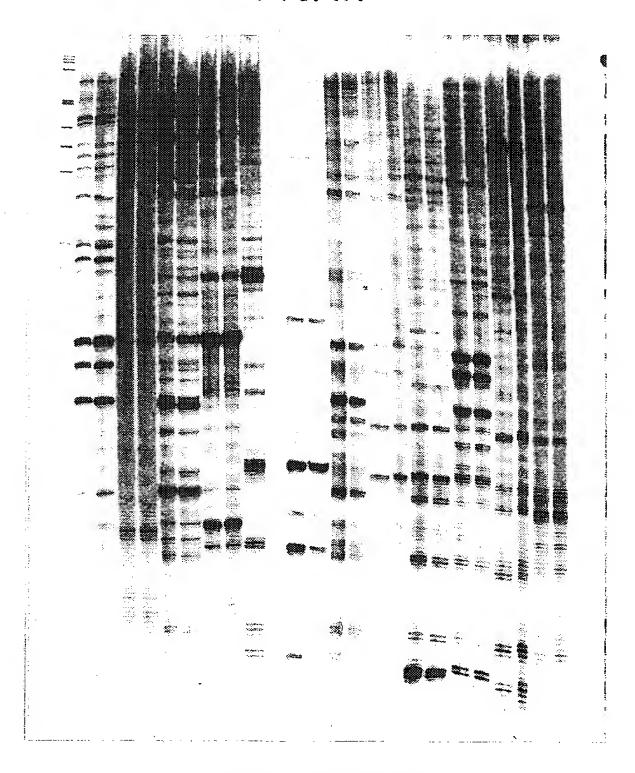
Methods are disclosed to identify T lymphocyte genes that are differentially expressed upon exposure to a pathogen (viral or bacterial), immunogen, antigen, or in a sterile inflammatory disease, autoimmune disease, immunodeficiency disease, lymphocytic cancers, or graft versus host rejection. The method involves the preparation of a gene expression profile of a T lymphocyte population exposed to a pathogen or isolated from a subject having one of the aforementioned pathologies and comparing that profile to a profile prepared from quiescent T lymphocytes. The present invention is particularly useful for identifying cytokine genes, genes encoding cell surface receptors and genes encoding intermediary signalling molecules. Related methods for identifying therapeutic or prophylactic immunomodulatory agents are presented. Articles of manufacture are disclosed that comprise selected grouping of nucleic acids, affixed to a solid support, that correspond to genes that are differentially expressed in various populations or subpopulations of T lymphocytes at variations stages of T cell differentiation, in quiescent versus activated T lymphocytes or normal versus diseased T lymphocytes.

# FOR THE PURPOSES OF INFORMATION ONLY

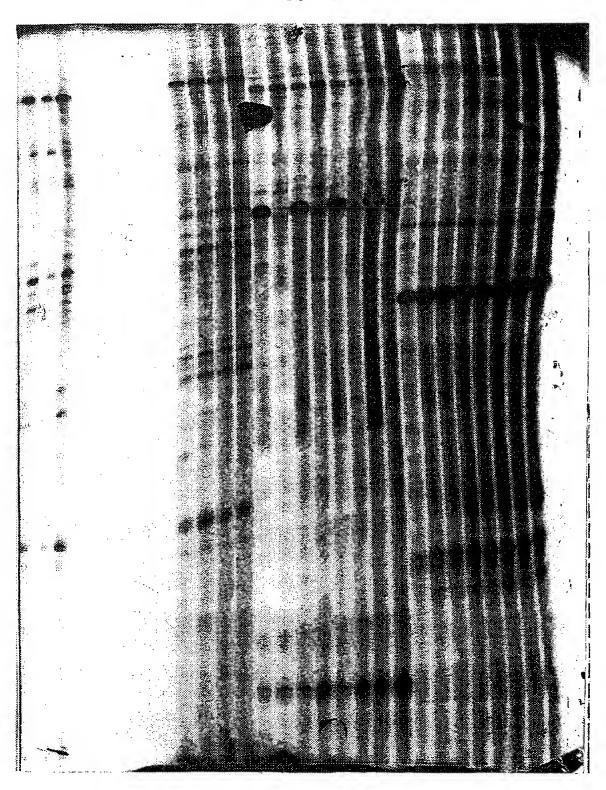
Codes used to identify States party to the PCT on the front pages of pamphlets publishing international applications under the PCT.

| AL | Albania                  | ES | Spain               | LS | Lesotho               | SI | Slovenia                 |
|----|--------------------------|----|---------------------|----|-----------------------|----|--------------------------|
| AM | Armenia                  | FI | Finland             | LT | Lithuania             | SK | Slovakia                 |
| AT | Austria                  | FR | France              | ŁU | Luxembourg            | SN | Senegal                  |
| ΑU | Australia                | GA | Gabon               | LV | Latvia                | SZ | Swaziland                |
| AZ | Azerbaijan               | GB | United Kingdom      | MC | Monaco                | TD | Chad                     |
| BA | Bosnia and Herzegovina   | GE | Georgia             | MD | Republic of Moldova   | TG | Togo                     |
| BB | Barbados                 | GH | Ghana               | MG | Madagascar            | TJ | Tajikistan               |
| BE | Belgium                  | GN | Guinea              | MK | The former Yugoslav   | TM | Turkmenislan             |
| BF | Burkina Faso             | GR | Greece              |    | Republic of Macedonia | TR | Turkey                   |
| BG | Bulgaria                 | HU | Hungary             | ML | Mali                  | TT | Trinidad and Tobago      |
| BJ | Benin                    | IE | Ireland             | MN | Mongolia              | UA | Ukraine                  |
| BR | Brazil                   | IL | Israel              | MR | Mauritania            | UG | Uganda                   |
| BY | Belarus                  | IS | Iceland             | MW | Malawi                | US | United States of America |
| CA | Canada                   | IT | Italy               | MX | Mexico                | UZ | Uzbekistan               |
| CF | Central African Republic | JР | Japan               | NE | Niger                 | VN | Viet Nam                 |
| CG | Congo                    | KE | Kenya               | NL | Netherlands           | YU | Yugoslavia               |
| CH | Switzerland              | KG | Kyrgyzstan          | NO | Norway                | ZW | Zimbabwe                 |
| CI | Côte d'Ivoire            | KP | Democratic People's | NZ | New Zealand           |    |                          |
| CM | Cameroon                 |    | Republic of Korea   | PL | Poland                |    |                          |
| CN | China                    | KR | Republic of Korea   | PT | Portugal              |    |                          |
| CU | Cuba                     | KZ | Kazakstan           | RO | Romania               |    |                          |
| CZ | Czech Republic           | LC | Saint Lucia         | RU | Russian Federation    |    |                          |
| DE | Germany                  | LI | Liechtenstein       | SD | Sudan                 |    |                          |
| DK | Denmark                  | LK | Sri Lanka           | SE | Sweden                |    |                          |
| EE | Estonia                  | LR | Liberia             | SG | Singapore             |    |                          |
|    |                          |    |                     |    |                       |    |                          |

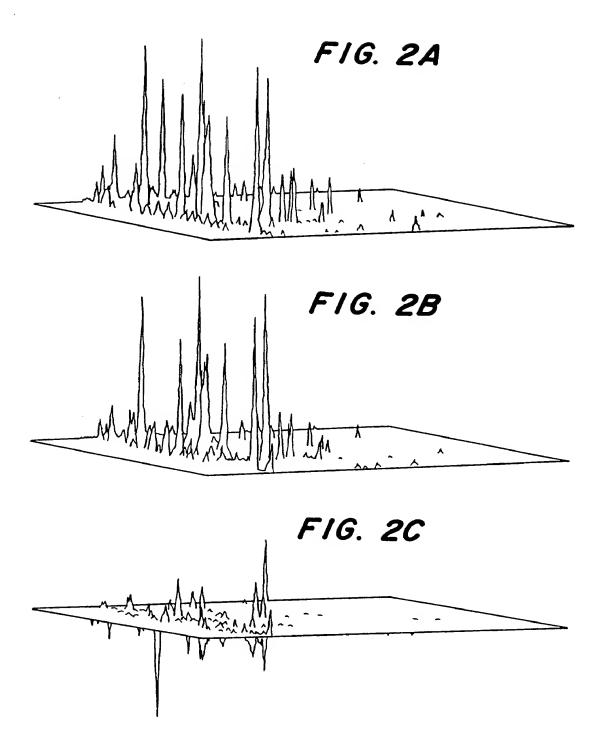
FIG. IA



# FIG. IB



SUBSTITUTE SHEET (RULE 26)



\* PARTIAL PATTERNS SHOWN REPRESENT APPROXIMATELY 5% OF COMPLETE PATTERN (ALL GENES EXPRESSED).

FIG. 3 TA1 1 8 JkA6 2 JkA1 JkA7 3 JkA2 JkA8 10 JkA3 11 JkA9 5 JkA4 JkA10 12 JkA5 6 JkA11 13 IL-2 JkR1 7(a) 15  $\beta\text{-Actin}$ 

SUBSTITUTE SHEET (RULE 26)

F16.4

|    |        |      |                     | Dil                 | Differential Display Summary of Results | ssults                        |
|----|--------|------|---------------------|---------------------|---|-------------------------------|
| Ö. | Clones | NINZ | Expression Patterns | n Patterns          | Gene Ba                                 | Gene Bank Search and Analysis |
| į  |        |      | Control T<br>cells  | Activated<br>Tcells |   |                               |
| -  | TAI    |      | 0                   | -1-                 | Unknown                                 |                               |
| CI | JkA1   |      | +                   | ‡                   | Unknown                                 |                               |
| 'n | JkA2   |      | -1-                 | ‡                   | Unknown                                 |                               |
| 4  | JkA3   |      | +/-                 | ‡                   | Unknown                                 |                               |
| 'n | JkA4   |      | -\+                 | #                   | Unknown                                 |                               |
| 9  | JkA5   |      | +                   | ‡                   | Unknown                                 |                               |
| 7  | JkA6   |      | 0                   | -+-                 | Unknown                                 |                               |
| ∞  | JkA7   |      | 0                   | ‡                   | Unknown                                 |                               |
| 6  | JkA8   |      | ‡                   | ‡                   | Unknown                                 |                               |
| 10 | JKA9   |      | ‡                   | ‡                   | Unknown                                 | -                             |
| 11 | JKA10  |      | +1-                 | ‡                   | Unknown                                 |                               |
| 12 | JKA11  |      | ‡                   | ‡                   | Human Serine Esterase                   |                               |
| 13 |        |      | 0                   | ‡                   |   |                               |
| 14 |        |      | 0                   | ‡                   | Human Ribosamal Protein 58              |                               |
|    |        |      |                     |                     |   |                               |

SUBSTITUTE SHEET (RULE 26)

F16. 4 cont.

|   |                               |                     | T -        |                                 |
|---|-------------------------------|---------------------|------------|---------------------------------|
| Differential Display Summary of Results | Gene Bank Search and Analysis |                     |            | n Factor SVJ1                   |
| ferential Display                       |                               |                     | ++ Unknown | Protein Translation Factor SVJ1 |
| Dif                                     | n Patterns                    | Activated<br>Tcells | ‡          | ‡                               |
|   | N1N2 Expression               | Control T           | 0          | 0                               |
|   | NIN                           |                     |            |                                 |
|   | No. Clones                    |                     |            |                                 |
|   | ė,                            |                     | 15         | 16                              |

# 7/8

| Clone v. Name                  | Enzyme | Band No. | R.P. | Level of<br>Regulation* | Match<br>(if any)                            | Percent ID | GenBank Accession | SEQ ID NO. |
|--------------------------------|--------|----------|------|-------------------------|--|------------|-------------------|------------|
| 02_SV_011398_S<br>AC7E_02      | Bln I  | 11       | 10.2 | 4X.1                    | Human mRNA for actin binding<br>protein p57  | 98.182     | HUMP57B           | 14         |
| 02_SV_011698_B<br>SSIBAND26_02 | Bln 1  | 4        | 8.5  | Absent                  | Human mRNA for KIAA0120 gene                 | 99.194     | HUMORFFA          | 15         |
| 04_SV102897_2A                 | Bln 1  | 12       | 10.2 | 2X 1                    | No human match                               |            |                   | 91         |
| 06_SV102897_1B                 | Bln I  | \$       | 8.5  | 1.5% i                  | Human ferritin light subunit mRNA            | 95.918     | HUMFERLA          | 17         |
| 06_SV_011398_S<br>AC14A_06     | Bln 1  | 13       | 10.2 | 2X I                    | Human STS WI-30177                           | 98.611     | G23064            | 18         |
| 08_SV102897_2B                 | Bln 1  | 9        | 9.4  | 2X 1                    |  |            |                   |            |
| 10_NS108E_10                   | Bln I  | 4        | 10.2 | 1.5X t                  | Human DNA sequence from BAC<br>397C4         | 94.554     | HS397C4           |            |
| 10_SV102897_IC                 | Bh I   | 7        | 9.4  | Absent                  | Human clone 23732 mRNA                       | 99.559     | HSU79258          | 20         |
| 10_SV_011398_S<br>AC15D_10     | Bln 1  | 15       | 10.2 | 3X i                    | Human hsc70 gene for 71 kd heat<br>shock     | 98.785     | HSHSC70           | 21         |
| 12_SV102897_2C                 | Bln l  | 18       | 10.3 | 1 X Z                   | Human colin carcinoma laminin-<br>binding pr | 99,602     | НОМLАМВ           |            |
| 14_SV102897_1D                 | Bln 1  | 6        | 9.4  | 2X I                    | No human match                               |            |                   | 22         |
| 16_SV102897_2D                 | Bln 1  | 10       | 9.4  | (Absent in R            | Human mRNA export protein Rael               | 98.788     | HSU84720          | 23         |
| 18_SV102897_1E                 | Nsi 1  | 80       | 8.4  | 2X1                     | No human match                               |            |                   | 24         |
| 19_SV011698_BS<br>SIBAND15_19  | Sac J  | -        | 8.3  | 2X1                     | No human match                               |            |                   | 25         |

# SUBSTITUTE SHEET (RULE 26)

F16. 5 cont.

| Clone v. Name                  | Enzyme | Band No. | R.P. | Level of<br>Regulation* | Match<br>(If any)                              | Percent ID | GenBank Accession | SEQ ID NO. |
|--------------------------------|--------|----------|------|-------------------------|--|------------|-------------------|------------|
| 20_SV102897_2E                 | Sac I  | 14       | 9.8  |                         | Human ferritin L chain mRNA                    | 100        | HUMFERL           | 26         |
| 23_SV011698_BS<br>S1BAND17_23  | Sac 1  | \$1      | 9.8  | 1.5X1                   | Human mRNA for Plede47                         | 100        | HUMP1CDC47        | 72         |
| 24_SV102897_2F                 | Sac I  | 7        | 8.4  | Absent                  | Human pancreatic rumor-related prot <b>ein</b> | 99.512     | HUMPANCAN         | 28         |
| 26_SV102897_1G                 | BssSl  | 15       | 8.6  | 2X1                     | Homo sapiens HLA-B gene                        | 100        | D83956            | 29         |
| 27_SV_011698_B<br>SSIBAND19_27 | BssSI  | 17       | 9.2  | 2X1                     | Homo sapiens MLN50 mRNA                        | 97.5       | HSMLN50           | 30         |
| 30_SV102897_1H                 | BssSl  | 19       | 9.4  | 2X1                     | No human match                                 |            |                   | 31         |
| 31_SV_011698_B<br>SSIBAND22_31 | BssSl  | 22       | 10.2 | 4X1                     | Human platlet activating factor<br>acetylh     | 97.826     | HSLIS10           | 32         |
| 37_SV_010898_S<br>AC_BAND1_37  | BssSl  | 26       | 10.3 | 2.5X1                   | No human match                                 |            |                   | 33         |

Gene expression in Activated tissue is represented as (1) or down (1) regulated with respect to Resting Jurkat tissue. X represents fold increase or decrease in expression levels.

SUBSTITUTE SHEET (RULE 26)

#### SEQUENCE LISTING

```
<110> Gene Logic, Inc., c/o Dr. Larry Tiffany
<120> Process to Study Changes in Gene Expression in T
     Lymphocytes
<130> gene logic 5004pr expression T lymphs
<140> 60/084,329
<141> 1998-05-05
<160> 44
<170> PatentIn Ver. 2.0
<210> 1
<211> 238
<212> DNA
<213> Jurkat cell cDNA
<400> 1
gatcctatgt nececcagg eggetggean theceaeggg aagtgteeac tgaggteect 60
gagatggata cctctacctg acatggcctg aagatgcagg gcagaggaat tqcccatqga 120
cagtgacgca aggactaggc tgggagggag cgtgccaacc ccttttgcct ctgggtttgg 180
ggagcggagg gcctcttctt ggtgccctgc ccccaataaa ggaactggac naaqagat 238
<210> 2
<211> 174
<212> DNA
<213> Jurkat cell cDNA
<400> 2
gatctcatga tgtggctgtt gggaagatgg tggggtttgt ttgccagctt ggagtcctat 60
taaatgaaag ccagcaactc atgttggtaa taggtctact gtgggaacag ttatccctaa 120
ccacagetca aaategetat catetttagn caaattaaaa tetatgtgge ageg
<210> 3
<211> 175
<212> DNA
<213> Jurkat cell cDNA
<400> 3
gatetggtga etggetttte gttetgtgtt ettggettee taaatttate tgeecatatg 60
catatgtttt gagaatttgt aaagtgagag acatgatcct attaaaataa gaagg
<210> 4
```

1

```
<211> 285
<212> DNA
<213> Jurkat cell cDNA
<400> 4
gatoctocat ggoocageaa ggoocaagat aaatoottna coaccoaggo accotgtgag 60
cccaacaggt taattagtcc attaatttta gtgggacctg catatgttga aaattaccaa 120
tactgactga catgtgatgc tgacctatga taaggttgag tatttattag atgggaaggg 180
aaatttgggg attatttatc ctcctgggga cagtttgggg aggattattt attgtattta 240
tattgaatta tgtacttttt tcaataaagt cttatttttg tggcg
                                                                  285
<210> 5
<211> 182
<212> DNA
<213> Jurkat cell cDNA
<400> 5
gatotgaaac coaggitagg catgacatti canceccaaa coctacetca tetginetga 60
aagacgctga aactnoctgg gatgttttcg ggnacaagaa tgtanatttn coctatooct 120
gnacttggtt taatcnaatc aatgtgtgta ttagaataaa agtcacagca tcccaaaagc 180
<210> 6
<211> 130
<212> DNA
<213> Jurkat cell cDNA
<400> 6
caggatetta aaaateecag eeatetaaat atgttteeca aeteeattaa gtaaggtaaa 60
ataatatttg tatttatgtt cagatgttga agctgtcatt ctcgaataaa actacacttt 120
                                                                   130
agaaatggcg
<210> 7
<211> 361
<212> DNA
<213> Jurkat cell cDNA
<400> 7
gatetttega ggceaggtge ceaggtettt cateaagage cecattteea agtgeteagt 60
ancceptttt ggccagtgen cececacea atgggacaag egcaggteca gtggeeteee 120
cagctgaccg caggcaggga acaaggcaga ccctagaggg ccaggccaca gcaggggctg 180
aggatgcctg gtgaatggat gcctgggaga atggatgcca gaattcacgc atgaggctct 240
gaacagggct gggaaaactt ccaaacgaag ggaagctcat gtcttggtgc actttgtgat 300
gatgcttcaa cagcaggact gagatgggga catttacaat aaacagaaat gtatgggctc 360
                                                                   361
<210> 8
<211> 176
```

```
<212> DNA
<213> Jurkat cell cDNA
<400> 8
ggatcttgca cgtatctgtt ttcctccccc atqaactaga aaaccactta ctcccagaat 60
tcaggtcgtg cttgttagta ctatatcacc aagtccattc atttaatgat ccaaaactgt 120
aatgttgcac tgtattccaa ataaagggta aaaacagaac caaagttata actccg
<210> 9
<211> 128
<212> DNA
<213> Jurkat cell cDNA
<400> 9
gatcaattct atgtctgact ttgaaattcc atttacaatg tagtatgttt tcaatgnnaa 60
accataaagt aacatccaag tgtttcatgg tttgttggga aggtaatttt aaaataaaac 120
aatttccg
                                                                   128
<210> 10
<211> 138
<212> DNA
<213> Jurkat cell cDNA
<400> 10
gatcaagtca ctgcatgttg agaagtatag gtataacttg tgaccatatc acagctcctt 60
tatttatgta gtttcttcac attttatgtg tacaatcaag catgcctgct gaccaaggcc 120
agaggtggag tggaagcg
                                                                   138
<210> 11
<211> 271
<212> DNA
<213> Jurkat cell cDNA
<400> 11
gatctcaaca ttgttggttt cttttgtttt tcatttggta caactttctt gaatttagaa 60
attacatett tgcagttetg ttaggtgete tgtaattaac etgaettata tgtgaacaat 120
tttcatgaga cagtcatttt taactaatga agtgattctt tctcactact atctgtattg 180
tqqaatqcac aaaattqtqt aqqtqctqaa tqctqtaaqq aqtttagqtt qtatqaattc 240
tacaacccta taataaattt tactctatac g
                                                                   271
<210> 12
<211> 186
<212> DNA
<213> Jurkat cell cDNA
<400> 12
gatccaaaac tatttgggan atgtatgggt agggtaaatc agtaagaggt gttatttgga 60
accttgtttt ggacagttta ccagttgcct tttatcccaa agttgttgta acctnctgtg 120
```

```
atacgatgct tcaagagaaa atgcggttat aaaaaatggt tcagaattaa acttctaatt 180
cattcg
<210> 13
<211> 171
<212> DNA
<213> Jurkat cell cDNA
<400> 13
ggatctgacc tccacggage cgctgtcccc gccccctgc tcccgtctgt ctgtcctgtc 60
attgctttga aaacatgact caataaaagt ttcctttcaa tttaaacacc g
<210> 14
<211> 151
<212> DNA
<213> T lymphocyte cDNA
<400> 14
agetecagaa gegettggae aggetggagg agacagteca ggecaagtag agececacag 60
ggcctccagc agggtcagcc attcacaccc atccactcac ctcccattcc cagccacgtg 120
gcagagaaaa aaatcataat aaaatggctt t
<210> 15
<211> 148
<212> DNA
<213> T lymphocyte cDNA
<400> 15
ttnngctacc tgngtccaag tcttggcttn ccctttccan tcacttcact gtgcgctaag 60
gggtggggtg aggggatgga gagggagggc tgcctaccat ggtctggggc ttgaggaaga 120
tgagtttgtt gatttaaata aagaattt
<210> 16
<211> 194
<212> DNA
<213> T lymphocyte cDNA
<400> 16
ctantttaga tacgtccana nccaggaccg ctgagaactg ggacagtttc ctgggatgag 60
tgccagcctg agcctgcatg gtgccgccga gcccggggtg gaggagggag ccaggcttcg 120
cttcaaggcg gcctctacct tttctcagaa tggtttcctg attgtgtcaa tgtgaaagtt 180
aaataaaatt tatg
<210> 17
<211> 116
<212> DNA
<213> T lymphocyte cDNA
```

```
<400> 17
cactginning aacggientg changianna nghettetge changinitet enetheance 60
aanaggcanc tttcntannt atcctaacaa gccttggacc aaatggaaat aacagc
<210> 18
<211> 212
<212> DNA
<213> T lymphocyte cDNA
<400> 18
getttattgg agagatacae acaaaggetg tecaeteaet tecataattt ettgatggae 60
atgtttttct cactgtcctt ctgcatgacc ttggctactg ccatctcaaa gtcctcctga 120
gtgacatgga ctcgccgttc tcgcagggca tacatgccag cttctgtgca cacgcccttc 180
acttcaagcc cctgatgctc ctggcatgag ct
                                                                   212
<210> 19
<211> 189
<212> DNA
<213> T lymphocyte cDNA
<400> 19
tgcatttatg gaaggcacat tacaggtctt tgtgggaaga aacagaaaga aatcacaaaa 60
gcaattaaga gagctcaaat aatggggttt atgccagtta catacaagga tcctgcatat 120
ctcaaggacc ctaaagtttg taacatcaga tatcgggaat aaattctatc acgttaccac 180
taataaact
                                                                   189
<210> 20
<211> 219
<212> DNA
<213> T lymphocyte cDNA
<400> 20
antgnaggga aagctatgaa aggtgccggc ggatctacaa catggaaatg gctcgcaaga 60
tcaacttctt gatgcgaaag aatcgggcag atccgtggca gggctgctga ggcctgtggg 120
tgggacaccc agtgcgaaac cctcatccag ttttctctcc atctctttc tttgtacaat 180
cccatttcct attaccattc tctgcaataa actcaaatc
                                                                   219
<210> 21
<211> 285
<212> DNA
<213> T lymphocyte cDNA
<400> 21
agetecteec tetggtggtg ettecteagg geceaceatt gaagaggttg attaageeaa 60
ccaagtgtag atgtagcatt gttccacaca tttaaaacat ttgaaggacc taaattcgta 120
gcaaattctg tggcagtttt aaaaagttaa gctgctatag taagttactg ggcattctca 180
atacttgaat atggaacata tgcacagggg aaggaaataa cattgcactt tataaacact 240
```

| gtattgtaag  | tggaaaatgc   | aatgtcttaa          | ataaaactat | ttaaa       |                          | 285 |
|-------------|--------------|---------------------|------------|-------------|--------------------------|-----|
| <210> 22    |              |                     |            |             |                          |     |
| <211> 195   |              |                     |            |             |                          |     |
| <212> DNA   |              |                     |            |             |                          |     |
| <213> T lyn | mphocyte cDM | NA .                |            |             |                          |     |
| -           | •            |                     |            |             |                          |     |
| <400> 22    |              |                     |            |             |                          |     |
| ctantttaga  | tncgtccaca   | gccaggaccg          | ctgagaactg | ggacagtttc  | ctgggatgag               | 60  |
| tgccagcctg  | agcctgcatg   | gtgccgccga          | gcccggggtg | gaggaggag   | ccaggcttcg               | 120 |
| cttcaaggcg  | gcctctacct   | tttctcagaa          | tggtttcctg | attgtgtcaa  | tgtgaaagtt               | 180 |
| aaataaaatt  | tatgt        |                     |            |             |                          | 195 |
|             |              |                     |            |             |                          |     |
| <210> 23    |              |                     |            |             |                          |     |
| <211> 180   |              |                     |            |             |                          |     |
| <212> DNA   |              |                     |            |             |                          |     |
| <213> T lyr | mphocyte cD1 | NA                  |            |             |                          |     |
| <400> 23    |              |                     |            |             |                          |     |
|             | ctctccattc   | cactacctat          | tacagaattt | ttetetaaaet | 2200000110               | 60  |
|             |              |                     |            |             | aagggggttg               |     |
|             |              |                     |            |             | ttcagtgtac<br>aataaagtct |     |
| gegeeagaga  | acaceggaaa   | agegeeege           | ageceegege | tgtatttgt   | aacaaagccc               | 100 |
| <210> 24    |              |                     |            |             |                          |     |
| <211> 138   |              |                     |            |             |                          |     |
| <212> DNA   |              |                     |            |             |                          |     |
| <213> T lyr | mphocyte cD1 | NA                  |            |             |                          |     |
|             |              |                     |            |             |                          |     |
| <400> 24    |              |                     |            |             |                          |     |
| aggntctctg  | agcacttacc   | gggcgtgacc          | gtttcttagg | tgtgagaggg  | gctgtggctt               | 60  |
| ttgtgcagcg  | actatgttgg   | tgttaggggt          | ggtgtggaga | ttgttaatct  | tgtataaagc               | 120 |
| aattcaataa  | attgtttc     |                     |            |             |                          | 138 |
|             |              |                     |            |             |                          |     |
| <210> 25    |              |                     |            |             |                          |     |
| <211> 74    |              |                     |            |             |                          |     |
| <212> DNA   |              |                     |            |             |                          |     |
| <213> T 1y  | mphocyte cD  | NA                  |            |             |                          |     |
| <400> 25    |              |                     |            |             |                          |     |
|             | ncttctgagg   | caatatatac          | acaagccttt | cagggggcac  | attcacaagt               | 60  |
| acctgttgtg  | _            | - 5 5 - 5 - 5 - 5 - |            | 53555       |                          | 74  |
|             |              |                     |            |             |                          | -   |
| <210> 26    |              |                     |            |             |                          |     |
| <211> 119   |              |                     |            |             |                          |     |
| <212> DNA   |              |                     |            |             |                          |     |
| <213> T ly  | mphocyte cD  | NA                  |            |             |                          |     |
|             |              |                     |            |             |                          |     |
| <400> 26    |              |                     |            |             |                          |     |

```
tgtntccntg naagggncct tgcanagtaa tagggcttct gcctaagcct ctccctccaa 60
gccaataggc agctttctta actatcctaa caagccttgg accaaatgga aataaagct 119
<210> 27
<211> 253
<212> DNA
<213> T lymphocyte cDNA
<400> 27
gtgnnccagt cttgncttgn ccaccgccca gnnacangct gntcngnatn antatgaaqa 60
gctcaatgtc tggcaggtca atgcttcccg gacacggatc acttttgtct gattccagcc 120
tgcttgcaac cctggggtcc tcttgttccc tgctggcctg ccccttggqa agggqcaqtq 180
atgcctttga ggggaaggag gagccctct ttctcccatg ctgcacttac tccttttgct 240
aataaaagtg ttt
<210> 28
<211> 344
<212> DNA
<213> T lymphocyte cDNA
<400> 28
cgagtgtagc acaancatnc gacnggcgac ttcgccantn tcatcctttn tgggaacanc 60
aanatacann ctccatttct ggagtcnggg tcttccgaag ccaggagctt gcctttccgc 120
tgagtccana ttggcaggtg gactacgagt Catacacatg gcggaaactg gatcctggca 180
gtgaggagac ccagacgctg gttcgagagt acttttcctg ggagggggcc ttccagcatg 240
tgggcaaagc cttcaatcag ggcaagatct tcaagtgaac atctcttgcc atcacctagc 300
tgcctgcacc tgcccttcag ggagatgggg gtcattaaag gaaa
<210> 29
<211> 456
<212> DNA
<213> T lymphocyte cDNA
<400> 29
agtgtntgcc cagggctctg atgtgtcnct canagcttgn nnagcctgac acagctgtct 60
tgtgagggac tgagatgcag gatttettea egecteneet ttgtgaette aanageetet 120
ggcatctett tetgcaaagg cacetgaatg tgtetgegte eetgttagen taatgtgagg 180
aggtggagag acageceace entgtgteea etgtgaeeee tgtteeeatg etgaeetgtg 240
tttcctcccc agtcatcttt cttgttccag agaggtgggg ctggatgtct ccatctctgt 300
ctcaacttta ngtgcactga gctgcaactt cttacttccc tactganaat aagaatctga 360
atatacattt gttttcccaa atatttggca tgaaaaggtt ntggataant taataagcca 420
ttcccgggat tttgggaaan caanttttac ctnnga
                                                                   456
<210> 30
<211> 122
<212> DNA
<213> T lymphocyte cDNA
```

```
<400> 30
cgtggngctc aagtettnan etgecenacg ggatcaaace tttenggeet gtnatgatte 60
tgaccatttg acttgannca cangtgaatc tttctcctgg tgactcaaat aaaagtataa 120
<210> 31
<211> 320
<212> DNA
<213> T lymphocyte cDNA
<400> 31
aggnanagtc catggggctg ccaacttcag acgaacagaa gaaacaggag attctgaaga 60
agttcatgga tcaacatccg gagatggatt tttccaaggc taaattcaac tagcccctgt 120
tttttcctcc ctgaactctt ggggctgagc tgcaaccacc caactttctt tcccactctt 180
ctctgggact tgtgggcctc agggcttggg gcaggcatgg gactggccca ggcacacagg 240
teceggggea teaggagaaa ggetgggtet tgggaeettg teeteecag ttggeetact 300
gttacacatt aaaacgattt
                                                                  320
<210> 32
<211> 116
<212> DNA
<213> T lymphocyte cDNA
gtgggtccaa gtctttgttt gnnctaagat ttgtnngctc tcagacngtg taaaacaaaa 60
tttattcatg ttttctgcat attaaaaaat cttattgtac caactggtaa actatt
<210> 33
<211> 210
<212> DNA
<213> T lymphocyte cDNA
<400> 33
tgtctccagg atctcatgag ccgcnacgtg ttnagagggt cnncatcata cgggggangg 60
ntggggcaaa tcgccacctg tacctttcct ctggccctgc tgcccccaca cccaactccg 120
anggeceacg etggggaaag egggaagege tegeteeett teececatta gtgetetete 180
tgcctggatc ccggcagaag ctatgaaagg
                                                                   210
<210> 34
<211> 155
<212> DNA
<213> T lymphocyte cDNA
<400> 34
tancntgnta cactegntaa agaagagean gateangena etataetana ngttageate 60
actaacgccc negcatgtgc atgaaacacc ttetetgene geenatteca natttacact 120
gggagaggtg ccagcaactg aataaatacc tctta
                                                                   155
```

| <210>      | 35               |                                       |    |
|------------|------------------|---------------------------------------|----|
| <211>      | 19               |                                       |    |
| <212>      | DNA              |                                       |    |
| <213>      | 5' sequence for  | primer                                |    |
| <400>      | 35               |                                       |    |
| ctctca     | aagga tctaccgct  |                                       | 19 |
| -2105      | 36               |                                       |    |
| <210><211> |                  |                                       |    |
| <211>      |                  |                                       |    |
|            |                  |                                       |    |
| (213)      | 5' sequence for  | primer                                |    |
| <400>      | 36               |                                       |    |
| cagggt     | agac gacgctacgc  |                                       | 20 |
| <210>      | 37               |                                       |    |
| <211>      | 20               |                                       |    |
| <212>      | DNA              |                                       |    |
| <213>      | 5' sequence for  | primer                                |    |
| -400       | 22               |                                       |    |
| <400>      | _                |                                       |    |
| taatad     | ccgcg ccacatagca |                                       | 20 |
| <210>      | 38               |                                       |    |
| <211>      | 55               |                                       |    |
| <212>      | DNA              |                                       |    |
| <213>      | 1-base anchored  | oligo(dT) primer                      |    |
| <400>      | 38               |                                       |    |
|            |                  | gggcgaattg ggtcgacttt ttttttttt ttttv | 55 |
|            |                  |                                       |    |
| <210>      |                  |                                       |    |
| <211>      |                  |                                       |    |
| <212>      |                  |                                       |    |
| <213>      | 2-base anchored  | oligo(dT) primer, RP5.0               |    |
| <400>      | 39               |                                       |    |
| ctctca     | agga tettaceget  | ttttttttt tttttttat                   | 40 |
| <210>      | 40               |                                       |    |
| <211>      | 40               |                                       |    |
| <212>      | DNA              |                                       |    |
| <213>      | 2-base anchored  | oligo(dT) primer, RP6.0               |    |
| <400>      | 40               |                                       |    |
|            |                  | ttttttttt tttttttcg                   | 40 |
|            |                  |                                       | 30 |

| <210> 41                                      |    |
|---|----|
| <211> 40                                      |    |
| <212> DNA                                     |    |
| <213> 2-base anchored oligo(dT) primer, RP9.2 |    |
|   |    |
| <400> 41                                      |    |
| cagggtagac gacgctacgc ttttttttt tttttttga     | 40 |
|   |    |
| <210> 42                                      |    |
| <211> 25                                      |    |
| <212> DNA                                     |    |
| <213> adapter oligonucleotide Al              |    |
|   |    |
| <400> 42                                      |    |
| tagcgtccgg cgcagcgacg gccag                   | 25 |
|   |    |
| <210> 43                                      |    |
| <211> 29                                      |    |
| <212> DNA                                     |    |
| <213> adapter oligonucleotide A2              |    |
| <400> 43                                      |    |
|   | 20 |
| gatcctggcc gtcggctgtc tgtcggcgc               | 29 |
| <210> 44                                      |    |
| <211> 40                                      |    |
| <212> DNA                                     |    |
| <213> PCR primer                              |    |
| <del>-</del>                                  |    |
| <400> 44                                      |    |
| tgaagccgag acgtcggtcg ttttttttt tttttttvn     | 40 |
|   |    |



From the INTERNATIONAL SEARCHING AUTHORITY

Facsimile No. (703) 305-3230

| To: REID G. ADLER MORGAN, LEWIS & BOCKIUS LLP 1800 M STREET, N.W.  | PCT  |  |  |  |
|--|--|--|--|--|
| WASHINGTON, D.C. 20036   | NOTIFICATION OF TRANSMITTAL OF<br>THE INTERNATIONAL SEARCH REPORT<br>OR THE DECLARATION  |  |  |  |
|  | (PCT Rule 44.1)  |  |  |  |
|  | Date of Mailing (day/month/year) 13 SEP 1999   |  |  |  |
| Applicant's or agent's file reference .  |  |  |  |  |
| 44921-5004WO   | FOR FURTHER ACTION See paragraphs 1 and 4 below  |  |  |  |
| International application No.  | International filing date  |  |  |  |
| PCT/US99/09761   | (day/month/year)<br>05 MAY 1999  |  |  |  |
| Applicant  | <del></del>  |  |  |  |
| GENE LOGIC, INC.   |  |  |  |  |
|  |  |  |  |  |
|  |  |  |  |  |
| 1. X The applicant is hereby notified that the internationa  | I search report has been established and is transmitted herewith.  |  |  |  |
| Filing of amendments and statement under Artic   | le 19:<br>the claims of the international application (see Rule 46):   |  |  |  |
| When? The time limit for liling such amendm  | ents is normally 2 months from the date of transmittal of the  |  |  |  |
| · ·  | more details, see the notes on the accompanying sheet.   |  |  |  |
| Where? Directly to the International Bureau of W<br>34, chemin des Colombe<br>1211 Geneva 20, Switzer<br>Facsimile No.: (41-22) 7  | ttes<br>rland  |  |  |  |
| For more detailed instructions, see the notes on   | the accompanying sheet.  |  |  |  |
| 2. The applicant is hereby notified that no internationa Article 17(2)(a) to that effect is transmitted herewith.  | I search report will be established and that the declaration under   |  |  |  |
| 3. With regard to the protest against payment of (an)  | additional fee(s) under Rule 40.2, the applicant is notified that:   |  |  |  |
| the protest together with the decision thereon has been transmitted to the International Bureau together with the applicant's request to forward the texts of both the protest and the decision thereon to the designated Offices. |  |  |  |  |
| no decision has been made yet on the protest; the applicant will be notified as soon as a decision is made.  |  |  |  |  |
| 4. Further action(s): The applicant is reminded of the fol   | lowing:  |  |  |  |
| the applicant wishes to avoid or postpone publication,   | ional application will be published by the International Bureau. If a notice of withdrawal of the international application, or of the provided in rules 90 bis 1 and 90 bis 3, respectively, before the al publication. |  |  |  |
|  | ternational preliminary examination must be filed if the applicant till 30 months from the priority date (in some Offices even later).   |  |  |  |
| Within 20 months from the priority date, the applicant must  | perform the prescribed acts for entry into the national phase before the demand or in a later election within 19 months from the priority  |  |  |  |
| Name and mailing address of the ISA/US   | Authorized of lice   |  |  |  |
| Commissioner of Patents and Trademarks Box PCT Washington, D.C., 20231   | PATRICK J. NOLAN   |  |  |  |

Telephone No.

(703) 308-0196



# **PCT**

## INTERNATIONAL SEARCH REPORT

(PCT Article 18 and Rules 43 and 44)

| Applicant's or agent's file reference<br>44921-5004WO | FOR FURTHER ACTION  | see Notification of<br>(Form PCT/ISA/220   | Transmittal of International Search Report ) as well as, where applicable, item 5 below.   |
|---|---|--|--|
| International application No.<br>PCT/US99/09761       | International filing date 05 MAY 1999   | (day/month/year)   | (Earliest) Priority Date (day/month/year) 05 MAY 1998                                      |
| Applicant<br>GENE LOGIC, INC.                         |   |  |  |
| This international search report consi                | ing transmitted to the Intern   | ational Bureau   | hority and is transmitted to the applicant   |
| 1. Certain claims were foun                           | d unsearchable (See Box I   | ).   | ·  |
| 2. Unity of invention is lack                         | ing (See Box II).   |  |  |
|   | rried out on the basis of the filed with the international furnished by the applicant but not acco                  | sequence listing application. separately from the ompanied by a statement the disclosure in the statement that the disclosure in the statement that the disclosure in the statement that | ent to the effect that it did not include matter<br>ne international application as filed. |
| 5. With regard to the abstract,                       |   | ed, according to Rule<br>may, within one n   | e 38.2(b), by this Authority as it appears nonth from the date of mailing of this          |
| 6. The figure of the drawings to be Figure No         | published with the abstract as suggested by the applicate because the applicant failer because this figure better c | ant.<br>d to suggest a figure  |  |



# INTERNATIONAL SEARCH REPORT

International application No.

|                       |  |                                   | PCT/US99/0976   | 1   |
|-----------------------|--|-----------------------------------|---|---|
| IPC(6) :              | SSIFICATION OF SUBJECT MATTER<br>(C07H 21/04; C12N 15/00; C12Q 1/02, 1/68)<br>(435/6, 29, 320.1, 325; 536/24.3, 23.5)<br>(International Patent Classification (IPC) or to both                           | national classification           | and IPC   |   |
| B. FIEL               | DS SEARCHED  |                                   |   |   |
| Minimum de            | ocumentation searched (classification system followe   | d by classification syn           | nbols)  |   |
| U.S. :                | 435/6, 29, 320.1, 325; 536/24.3, 23.5  |                                   |   |   |
| Documentat            | ion scarehed other than minimum documentation to the   | e extent that such docu           | ments are included                                    | in the fields searched  |
|                       | ata base consulted during the international search (na LOG MEDLINE EMBASE BIOSIS LIFESCI   | ame of data base and,             | where practicable,                                    | search terms used)  |
| C. DOC                | UMENTS CONSIDERED TO BE RELEVANT   |                                   |   |   |
| Category*             | Citation of document, with indication, where ap  | propriate, of the relev           | ant passages  | Relevant to claim No.   |
| Y                     | WO 97/05286 A1 (YALE UNIVERS entire document.  | ITY) 13 Februa                    | ry 1997, see  | 1-32  |
| Υ                     | GROSSMAN et al. SHORT ANALY Infection to AIDS: Are the Manifes Resistance Misinterpreted? Cli Immunopathology. 01 November 1993 135, see entire document.  | tations of Effec<br>nical Immuno  | tive Immune<br>ology And                              | 1-32  |
|                       | cr documents are listed in the continuation of Box C   |                                   | nt family annex.                                      |   |
| "A" doe               | ecial categories of cited documents:  cument defining the general state of the art which is not considered  be of particular relevance   | date and not i<br>the principle o | n conflict with the appli<br>or theory underlying the | rnational filing date or priority cation but cited to understand invention  claimed invention cannot be |
| *L* do:               | ther document published on or after the international filing date<br>cument which may throw doubts on priority claim(s) or which is<br>ed to establish the publication date of another citation or other | considered not<br>when the docu   | vel or camot be consider<br>ument is taken alone      | ed to involve an inventive step   |
| O" doe                | coal reason (as specified) cuntent referring to an oral disclosure, use, exhibition or other ans   | considered to combined with       | involve an inventive                                  | claimed invention earnot be<br>step when the document is<br>documents, such combination<br>be art       |
| the                   | cument published prior to the international filing date but later than<br>priority date claimed  | *&* document men                  | mber of the same patent                               | family  |
| Date of the           | actual completion of the international search  | Date of mailing of the 13 SEP     |   | rch report  |
| Commission<br>Box PCT | nailing address of the ISA/US<br>ner of Palents and Trademarks<br>n. D.C. 20231  | Australia de etta                 | aurexce   | - Tex   |
| Faccimile N           | 0 (703) 305,3330   | LT deshare No.                    | 1031 200 0106   |   |